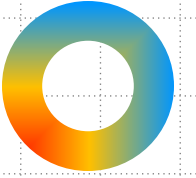




Python Programming

Principal Component Analysis

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Outlines

- Review
- Why Do We Need Dimension Reduction?
- PCA – Geometry
- PCA – Linear Algebra
- PCA – Characteristics
- PCA – Principal Components
- PCA – Variable Loading
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- PCA – sklearn.
decomposition.pca
- PCA – SVD
- PCA – Eigen
Decomposition
- PCA– Variable Loading

Review

- Before we explain PCA, we need to review the mathematical meaning of three basic descriptive statistics, including expectation, variance, and covariance.
- In previous courses or your understanding, these three parameters usually perform as above equations.

$$\mu = \frac{1}{n} \sum_{i=1}^n x_i$$

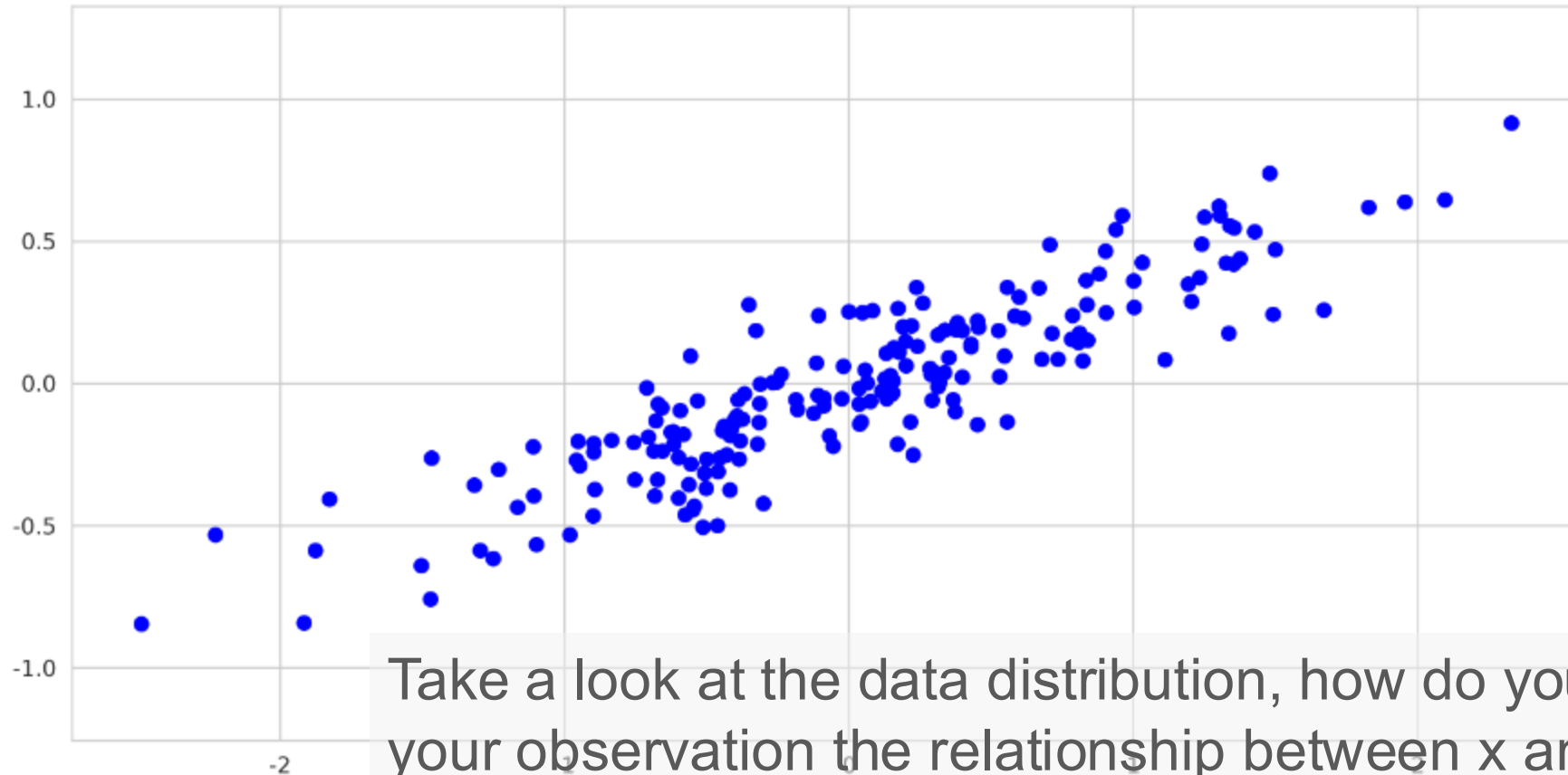
$$var(x) = \sigma^2 = \left(\sqrt{\frac{\sum_{i=1}^n (x_i - \mu)^2}{n}} \right)^2$$

$$= \frac{1}{n} \sum_{i=1}^n (x_i - \mu)^2$$

$$Cov(x, y) = \frac{1}{n} \sum_{i=1}^n (x_i - \mu_x)(y_i - \mu_y)$$

Review

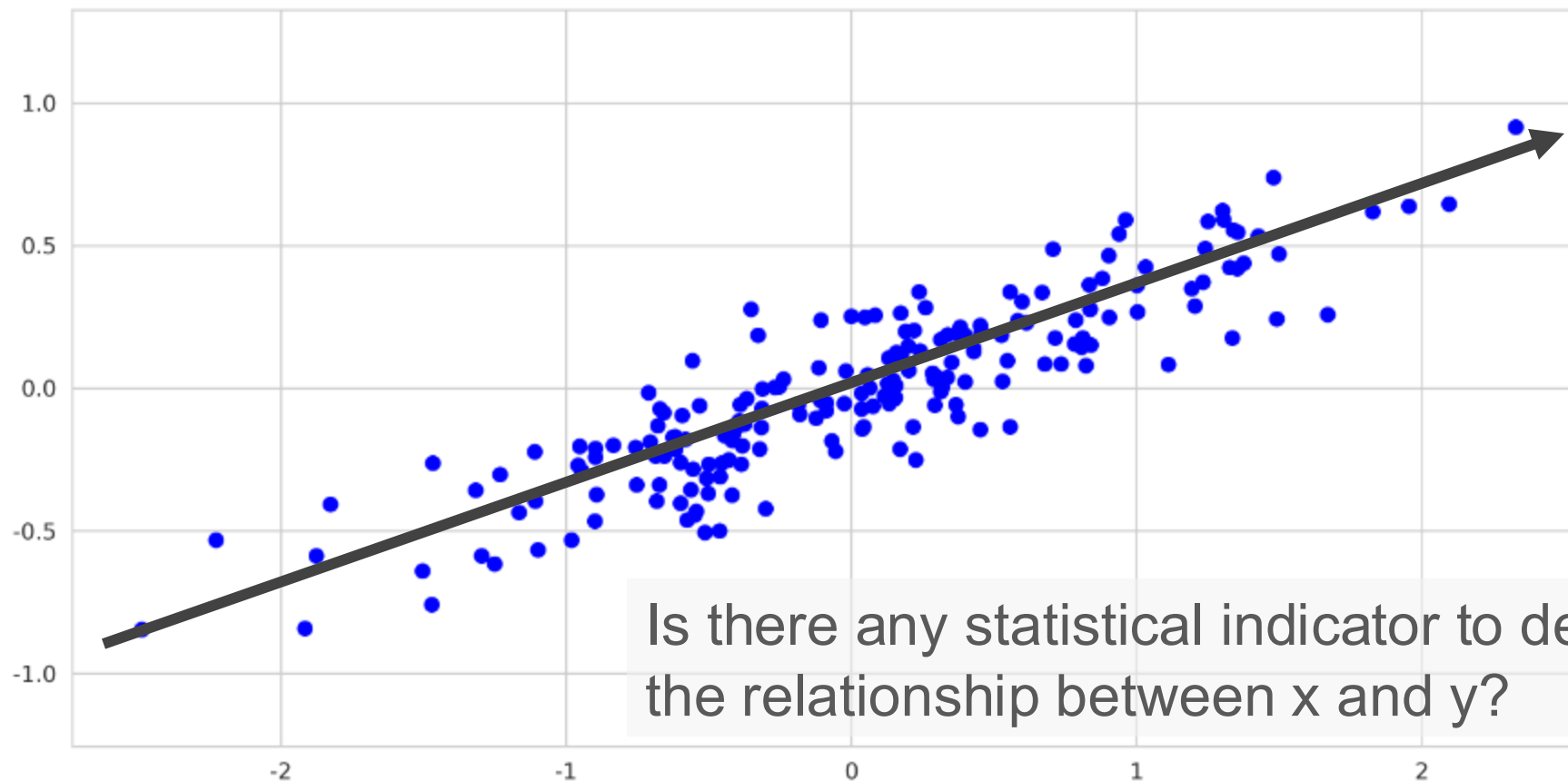
$$\mu = \frac{1}{n} \sum_{i=1}^n x_i \mid \text{var}(x) = \sigma^2 = \frac{1}{n} \sum_{i=1}^n (x_i - \mu)^2 \mid \text{Cov}(x, y) = \frac{1}{n} \sum_{i=1}^n (x_i - \mu_x)(y_i - \mu_y)$$



Take a look at the data distribution, how do you explain your observation the relationship between x and y?

Review

$$\mu = \frac{1}{n} \sum_{i=1}^n x_i \mid \text{var}(x) = \sigma^2 = \frac{1}{n} \sum_{i=1}^n (x_i - \mu)^2 \mid \text{Cov}(x, y) = \frac{1}{n} \sum_{i=1}^n (x_i - \mu_x)(y_i - \mu_y)$$

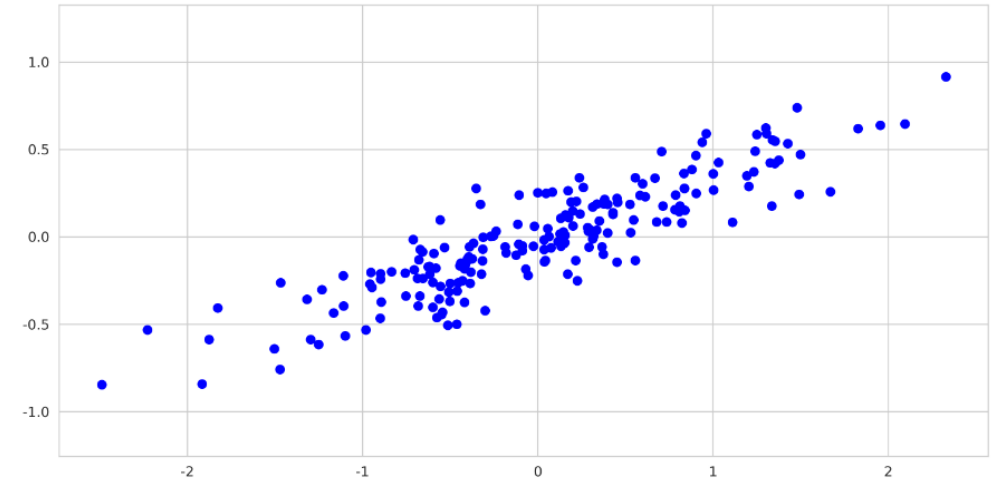


Review

$$\mu = \frac{1}{n} \sum_{i=1}^n x_i \mid \text{var}(x) = \sigma^2 = \frac{1}{n} \sum_{i=1}^n (x_i - \mu)^2 \mid \text{Cov}(x, y) = \frac{1}{n} \sum_{i=1}^n (x_i - \mu_x)(y_i - \mu_y)$$

- Pearson's correlation coefficient
- Given two parameters x_i and y_i , where i ranges from 1 to n . Then, Pearson's correlation coefficient could be defined as follows.

$$\rho = \frac{\sum_{i=1}^n (x_i - \mu_x)(y_i - \mu_y)}{\sqrt{\sum_{i=1}^n (x_i - \mu_x)^2 \sum_{i=1}^n (y_i - \mu_y)^2}}$$



Question 1

If x is highly correlated with y , and then what do you expect from their covariance and standard deviations?

Review

$$\mu = \frac{1}{n} \sum_{i=1}^n x_i \mid \text{var}(x) = \sigma^2 = \frac{1}{n} \sum_{i=1}^n (x_i - \mu)^2 \mid \text{Cov}(x, y) = \frac{1}{n} \sum_{i=1}^n (x_i - \mu_x)(y_i - \mu_y)$$

- **Expectation**

$$E(X) = \sum_x xP(X = x) = \mu$$

- **Variance**

$$\begin{aligned} \text{var}(X) &= E([X - \mu]^2) \\ &= E(X^2 - 2\mu X + \mu^2) \\ &= E(X^2) - 2\mu E(X) + \mu^2 \\ &= E(X^2) - 2\mu^2 + \mu^2 \\ &= E(X^2) - \mu^2 \\ &= E(X^2) - E(X)^2 \end{aligned}$$

Characteristics of Expectation

$E(aX + bY) = aE(X) + bE(Y)$, $a, b \in \mathbb{R}$
 X and Y can be independent or dependent.

$$E(XY) = E(X)E(Y)$$

Where $\text{cov}(X, Y) = 0$

Review

Covariance

$$E(aX + bY) = aE(X) + bE(Y), a, b \in \mathbb{R}$$

X and Y can be independent or dependent.

$$E(XY) = E(X)E(Y)$$

Where $cov(X, Y) = 0$

- If x and y are independent...

$$var(X + Y) = var(X) + var(Y)$$

- If x and y are dependent...

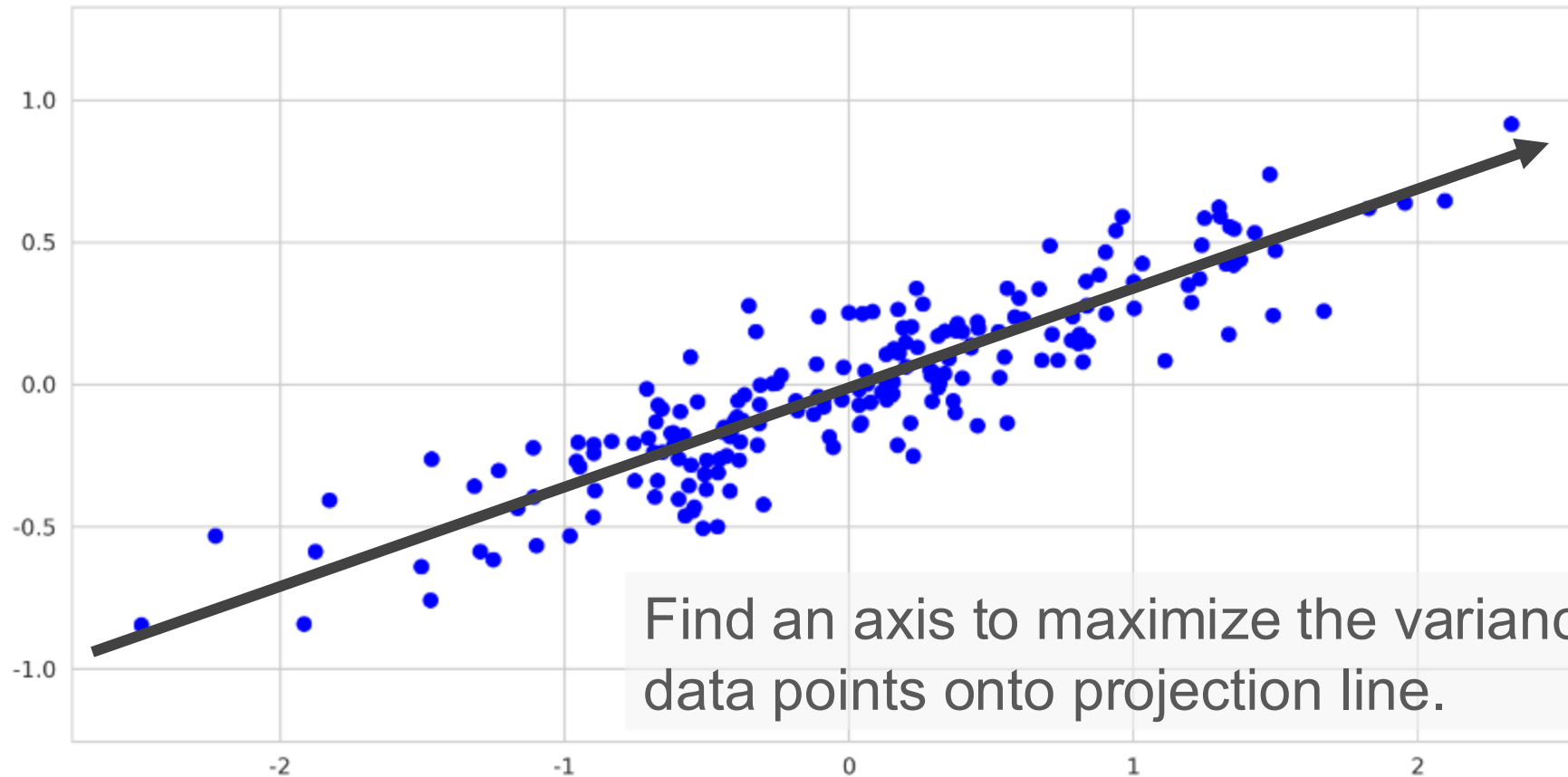
$$\begin{aligned} var(X + Y) &= E([(X + Y) - E(X + Y)]^2) \\ &= E\left([(X + Y) - (E(X) + E(Y))]\right)^2 \\ &= E\left([(X - E(X)) + (Y - E(Y))]\right)^2 \\ &= E\left((X - E(X))^2 + 2(X - E(X))(Y - E(Y)) + (Y - E(Y))^2\right) \\ &= E\left[(X - E(X))^2\right] + E\left[(Y - E(Y))^2\right] + 2E[(X - E(X))(Y - E(Y))] \\ &= var(X) + var(Y) + 2cov(X, Y) \end{aligned}$$

Why Do We Need Dimension Reduction?

- Here comes the first question into your mind.
 - Why do we need dimension reduction?
 - What's the importance of dimension reduction?
 - Can we directly import all datasets into your model without dimension reduction?
- Statistical models (e.g., linear regression) have several assumptions when you adopt them. One of them is “all variables have to be linearly independent,” indicating no collinearity.
- To achieve this goal, various methods were developed for orthogonalizing parameters and reducing the dimension of the dataset, such as PCA, LDA, LLE, and Laplacian Eigenmaps.

PCA – Geometry

$$\mu = \frac{1}{n} \sum_{i=1}^n x_i \quad | \quad \text{var}(x) = \sigma^2 = \frac{1}{n} \sum_{i=1}^n (x_i - \mu)^2 \quad | \quad \text{Cov}(x, y) = \frac{1}{n} \sum_{i=1}^n (x_i - \mu_x)(y_i - \mu_y)$$

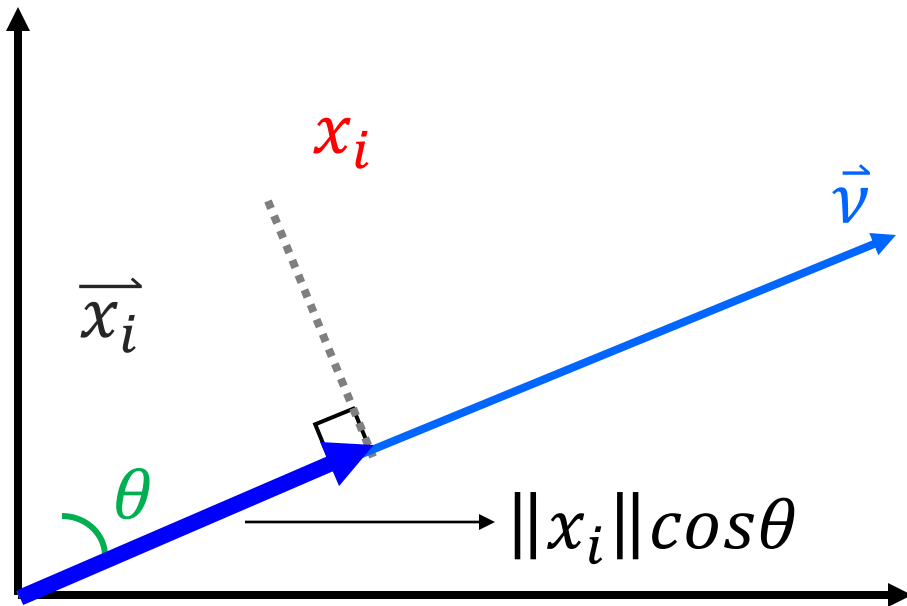


Find an axis to maximize the variance of all data points onto projection line.

PCA – Geometry

- Given a point “ x ” and project onto a vector “ v ”.

$$\cos\theta = \frac{x_i^T \cdot v}{\|x_i\| \|v\|}$$



$$\|x_i\| \cos\theta = \|x_i\| \frac{x_i^T \cdot v}{\|x_i\| \|v\|} = \frac{x_i^T \cdot v}{\|v\|}$$

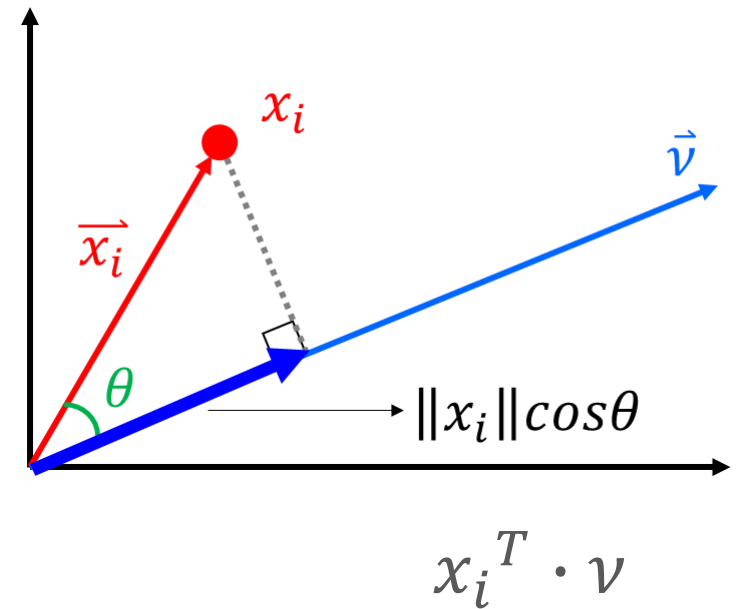
if v is unit vector ... $\|v\| = 1$

$$= \frac{x_i^T \cdot v}{\|v\|} = x_i^T \cdot v$$

PCA – Linear Algebra

$$X = \begin{bmatrix} | & | & \cdots & | \\ x_1 & x_2 & \cdots & x_n \\ | & | & \cdots & | \end{bmatrix} \rightarrow X^T = \begin{bmatrix} - & x_1 & - \\ - & x_2 & - \\ - & \vdots & - \\ - & x_n & - \end{bmatrix}$$

$$P = \begin{bmatrix} x_1^T \\ x_2^T \\ \vdots \\ x_n^T \end{bmatrix} = X^T u \Rightarrow \text{solve } P$$



PCA – Linear Algebra

$$J(u) = \|P^2\| = P^T P = (X^T u)^T (X^T u) = u^T X X^T u$$
$$\operatorname{argmax}_u J(u) = u^T X X^T u, \text{ where subject to } u^T u = 1$$

Add Lagrange multiplier

$$\operatorname{argmax}_{u, \lambda} J(u, \lambda) = u^T X X^T u + \lambda(1 - u^T u)$$

$$\nabla_u J(u, \lambda) = \nabla_u (u^T X X^T u + \lambda(1 - u^T u)) = 0$$

$$\Rightarrow 2X X^T u - 2\lambda u = 0$$

$$\Rightarrow \boxed{X X^T} u = \boxed{\lambda} u \rightarrow \text{eigenvector} \xrightarrow{\text{yellow arrow}} Au = \lambda u$$

\downarrow \downarrow
 $\text{cov}(X)$ eigenvalue

PCA – Linear Algebra

When u is the eigenvector

$$J(u) = \|P^2\| = u^T X X^T u = u^T \lambda u = \lambda u^T u = \lambda$$

$$X X^T u = \lambda u$$

*Given an **eigenvector**, the **total square of projected values** is the **eigenvalue** $= \lambda$*

An eigenvector is a symmetry matrix
 u is a unit vector

$$u u^T = u^T u = 1$$

PCA – Characteristics

$$XX^T u = \lambda u$$

A is a square symmetric matrix has orthogonal eigenvectors with different eigenvalues.

An eigenvector is a symmetry matrix
 u is a unit vector

$$uu^T = u^T u = 1$$

$$[x_1, \lambda_1], [x_2, \lambda_2]$$

$$\begin{cases} Ax_1 = \lambda_1 x_1 \\ Ax_2 = \lambda_2 x_2 \end{cases}$$

$$x_1^T Ax_2 = x_1^T \lambda_2 x_2 = \lambda_2 x_1^T x_2$$

Equal

$$x_1^T A^T x_2 = (Ax_1)^T x_2 = (\lambda_1 x_1)^T x_2 = \lambda_1 x_1^T x_2$$

$$(\because A \in \text{symmetric matrix}, \therefore A = A^T)$$

$$\begin{aligned} \lambda_2 x_1^T x_2 &= \lambda_1 x_1^T x_2 \\ \underline{x_1^T x_2} (\lambda_2 - \lambda_1) &= 0 \end{aligned}$$

Orthogonal $x_1^T x_2 = 0$ All eigenvalues are different

PCA – Characteristics

- Conversion between orthogonal bases

$$u_i \cdot u_j = u_i^T \cdot u_j = \begin{cases} 1, & \text{if } i = j \\ 0, & \text{otherwise} \end{cases}$$

$$U = \begin{bmatrix} | & | & & | \\ u_1 & u_2 & \dots & u_d \\ | & | & & | \end{bmatrix} \Rightarrow U^T U = I = U^{-1} = U^T$$

$$x = y_1 u_1 + y_2 u_2 + \dots + y_d u_d = \begin{bmatrix} | & | & & | \\ u_1 & u_2 & \dots & u_d \\ | & | & & | \end{bmatrix} \begin{bmatrix} y_1 \\ y_2 \\ \vdots \\ y_d \end{bmatrix} = Uy$$

$$\Rightarrow y = U^{-1}x = U^T x$$

PCA – Princial Components

Principal components (PC)

- Centered Matrix ($A^{N \times p}$ with N samples and p features)

$$A_{centered} = A - \bar{A}$$

- Compute the covariance matrix

$$S = \frac{1}{N} A_{centered}^T A_{centered}, S \in p \times p \text{ matrix}$$

- Apply Eigen Decomposition

$$SV = \lambda V$$

V (eigenvectors) define **principal component directions**

λ (eigenvalues) define **variance explained by each PC**

PCA – Principal Components

- The eigenvectors of S are the Principal Components (PC s)

$$PC = A_{centered}V$$

- Using Singular Value Decomposition (SVD)

$$A_{centered} = U\Sigma V^T$$

U contains left singular vectors

Σ contains singular values

V contains right singular vectors (principal component directions)

$$PC \Rightarrow A_{centered}V = U\Sigma V^TV$$

$$\because V^TV = I, \therefore \mathbf{A_{centered}V = U\Sigma = PC}$$

*Each **row** of PC is a transformed data point*

***Columns** of PC correspond to principal components.*

PCA – Principal Components

- **Compute PCA to Covariance Matrix**
- PCA is based on the Eigen Decomposition of the covariance matrix

$$S = \frac{1}{N} A^T A$$

- Using Singular Value Decomposition (SVD)

$$A^T A = (U \Sigma V^T)^T (U \Sigma V^T) = V \Sigma^T U^T U \Sigma V^T$$

- Since U is orthogonal, $U^T U = I$

$$A^T A = V \Sigma^T \Sigma V^T$$

- Divide by N

$$\frac{1}{N} A^T A = \frac{1}{N} V \Sigma^T \Sigma V^T$$

PCA – Principal Components

$$\frac{1}{N} A^T A = \frac{1}{N} V \Sigma^T \Sigma V^T = V \frac{\Sigma^T \Sigma}{N} V^T$$

- From Eigen Decomposition, the **eigenvalues** of S are:

$$E = \frac{\Sigma^T \Sigma}{N}$$

- Taking the square root:

$$\Sigma = \sqrt{N} \sqrt{E}$$

PCA – Variable Loading

- From the covariance matrix, we can obtain variable loading

$$\text{cov}(A, PC) = \frac{1}{N} A^T PC, \because \text{cov}(X, Y) = \mathbb{E}[(X - \mu_x)(Y - \mu_y)]$$

- Substitution with $PC = U\Sigma = A_{\text{centered}}V$

$$\text{cov}(A, PC) = \frac{1}{N} A^T U \Sigma, \text{ where } \Sigma = \sqrt{NE}$$

- Since E is just a scaling factor; therefore, we may approximate

$$\text{cov}(A, PC) = \frac{1}{N} A^T U \sqrt{N} = \frac{\sqrt{N}}{N} A^T U = \frac{1}{\sqrt{N}} A^T U$$

PCA – Variable Loading

$$\text{cov}(A, PC) = \frac{1}{\sqrt{N}} A^T U$$

- Since Singular Value Decomposition (SVD)

$$A^T = (U \Sigma V^T)^T = V \Sigma^T U^T$$

$$\text{cov}(A, PC) = \frac{1}{\sqrt{N}} V \Sigma^T U^T U$$

- Since U is an orthogonal matrix, $\therefore U^T U = I$, and Σ is a diagonal matrix with singular values, $\therefore \Sigma^T = \Sigma$, and $\Sigma = \sqrt{NE}$

$$\text{cov}(A, PC) = \frac{1}{\sqrt{N}} V \Sigma = V \frac{1}{\sqrt{N}} \sqrt{NE} = V \sqrt{E}$$

PCA – Variable Loading

- Finally, the variable loading could be obtained

$$\text{cov}(A, PC) = \mathcal{L} = V\sqrt{E}$$

\mathcal{L} is the variable loading of each PC

V is the eigenvector of A

E is the eigenvalues of A

PCA – Steps of PCA

The Steps of PCA

1. Find the sample mean $\mu = \frac{1}{n} \sum_{i=1}^n x_i$
2. Subtract mean
3. Compute covariance matrix $C = \frac{1}{n} X X^T = \frac{1}{n} \sum_{i=1}^n (x_i - \mu) (x_i - \mu)$
4. Find the eigenvalues of C and arrange them into descending order
 $\lambda_1 > \lambda_2 > \dots > \lambda_d, \{u_1, u_2, \dots, u_d\}$
5. The transformation is $y = U^T X$.

PCA – Python Tutorial

- As mentioned earlier, Principal Component Analysis (PCA) can be computed using different methods, each with its own advantages:

1) Pre-built Libraries:

The simplest approach is using `sklearn.decomposition.PCA`, which internally applies Singular Value Decomposition (SVD) for numerical stability and efficiency.

PCA – Python Tutorial

2) Eigen Decomposition:

This method computes eigenvalues and eigenvectors of the covariance matrix ($A^T A / N$), where eigenvectors define the principal component directions and eigenvalues indicate the explained variance. It works best when the covariance matrix is well-conditioned.

3) Singular Value Decomposition (SVD):

Instead of explicitly computing the covariance matrix, SVD decomposes the mean-centered data matrix as $A = U\Sigma V^T$.

- While eigen decomposition offers an intuitive understanding, SVD is generally preferred for its robustness and efficiency.

PCA – sklearn.decomposition.pca

define a matrix with 200 samples with two features

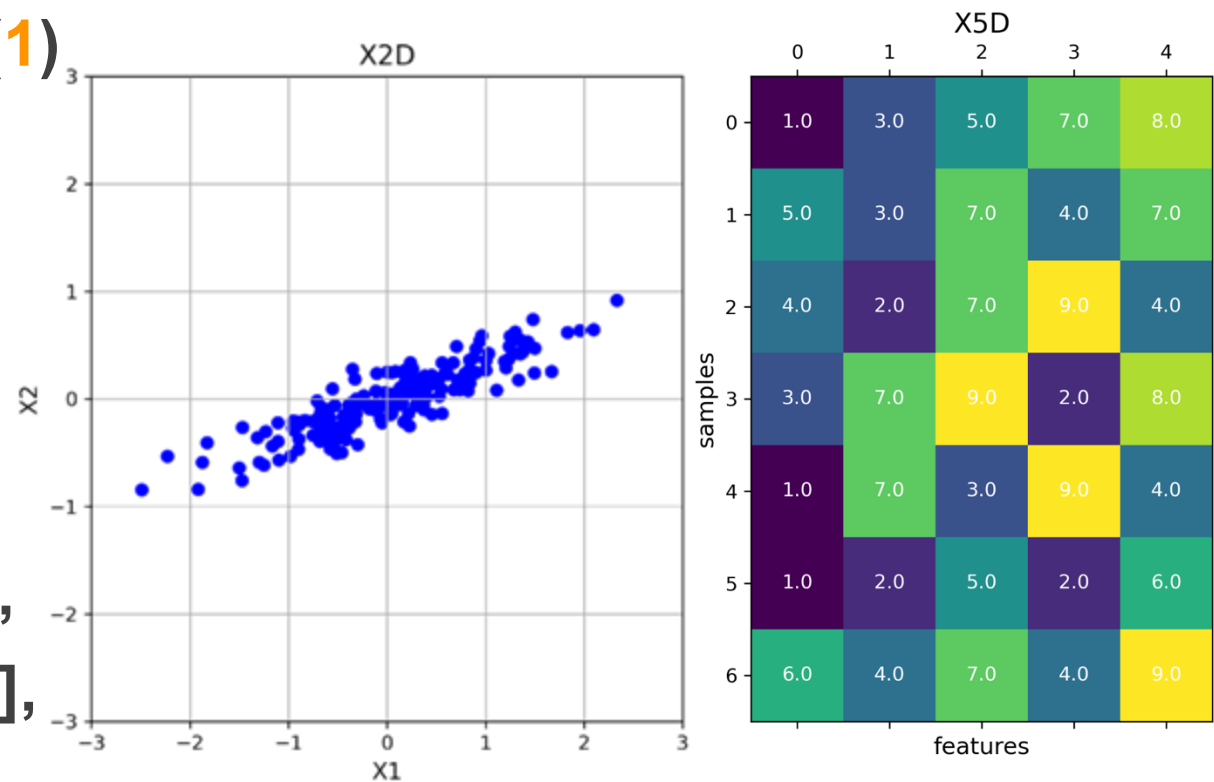
```
rn_state = np.random.RandomState(1)
```

2x200 -> 200x2

```
A2D = np.dot(rn_state.rand(2, 2),  
             rn_state.randn(2, 200)).T
```

define a matrix with seven samples
and five features

```
A5D = np.array([[1, 3, 5, 7, 8], [5, 3, 7, 4, 7],  
                [4, 2, 7, 9, 4], [3, 7, 9, 2, 8], [1, 7, 3, 9, 4],  
                [1, 2, 5, 2, 6], [6, 4, 7, 4, 9]])
```



PCA – sklearn.decomposition.pca

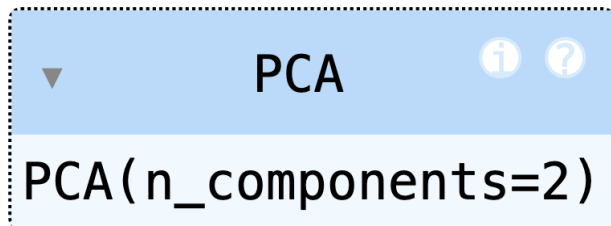
set the number of components for PCA model

```
pca2D = PCA(n_components=2)
```

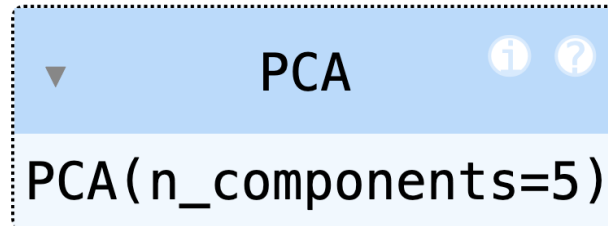
```
pca5D = PCA(n_components=5)
```

fit PCA model

```
pca2D.fit(A2D)
```



```
pca5D.fit(A5D)
```



PCA – sklearn.decomposition.pca

```
# principal component - eigenvector
```

```
# (n_components, n_features)
```

```
print(pca2D.components_)
```

```
[[-0.94446029 -0.32862557]  
 [-0.32862557  0.94446029]]
```

```
print(pca5D.components_)
```

```
[[-2.84777513e-01  1.33828752e-02 -4.01328150e-01  7.56452377e-01  
 -4.30625340e-01]  
 [ 6.37290518e-01 -6.01836596e-01  2.90984089e-01  3.81084174e-01  
 -4.19120474e-02]  
 [-4.04144877e-01 -7.88343623e-01 -3.10944893e-01 -3.38567133e-01  
 -6.21837647e-02]  
 [-4.55042776e-04 -5.10904099e-02 -3.73205245e-01  3.01380387e-01  
  8.75943646e-01]  
 [-5.91125431e-01 -1.16265701e-01  7.19927145e-01  2.77663127e-01  
  2.04110524e-01]]
```

PCA – sklearn.decomposition.pca

```
# eigenvalues
```

```
print(pca2D.explained_variance_)
```

```
print(pca5D.explained_variance_)
```

```
[0.7625315 0.0184779]
```

```
[13.48994148  5.38421307  4.33804606  1.80323359  0.98456579]
```

```
# variance explained ratio
```

```
print(pca2D.explained_variance_ratio_)
```

```
print(pca5D.explained_variance_ratio_)
```

```
[0.97634101 0.02365899]
```

```
[0.5188439  0.20708512 0.16684793 0.06935514 0.03786791]
```

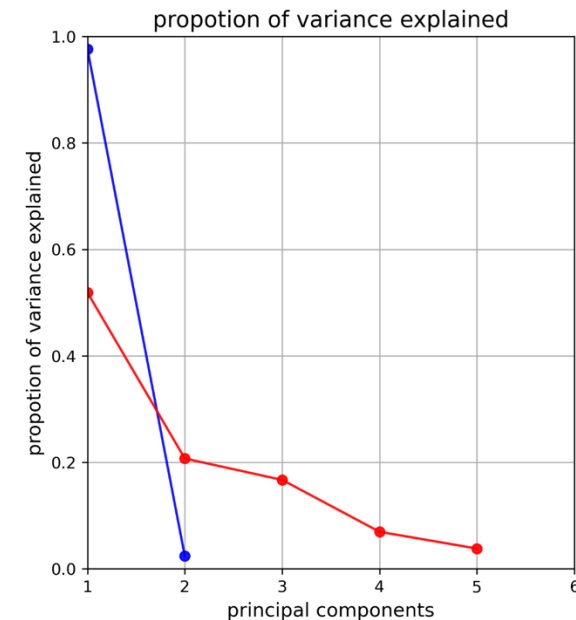
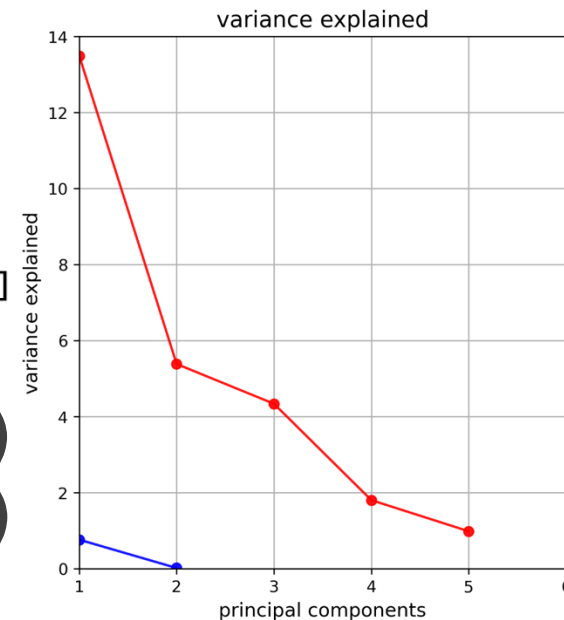
```
# variance explained ratio
```

```
print(pca2D.explained_variance_/sum(pca2D.explained_variance_))
```

```
print(pca5D.explained_variance_/sum(pca5D.explained_variance_))
```

```
[0.97634101 0.02365899]
```

```
[0.5188439  0.20708512 0.16684793 0.06935514 0.03786791]
```



PCA – sklearn.decomposition.pca

```
# project data onto PC1
```

```
pca1D = PCA(n_components=1)
```

```
pca1D.fit(A2D)
```

```
#  $A_{pca} = A_{centered} \times v$ 
```

```
A2D_pca1D = pca1D.transform(A2D)
```

```
A5D_pca5D = pca5D.transform(A5D)
```

```
print("original shape: ", A2D.shape) original shape: (200, 2)
```

```
print("transformed shape:", A2D_pca1D.shape) transformed shape: (200, 1)
```

```
print("original shape: ", A5D.shape) original shape: (7, 5)
```

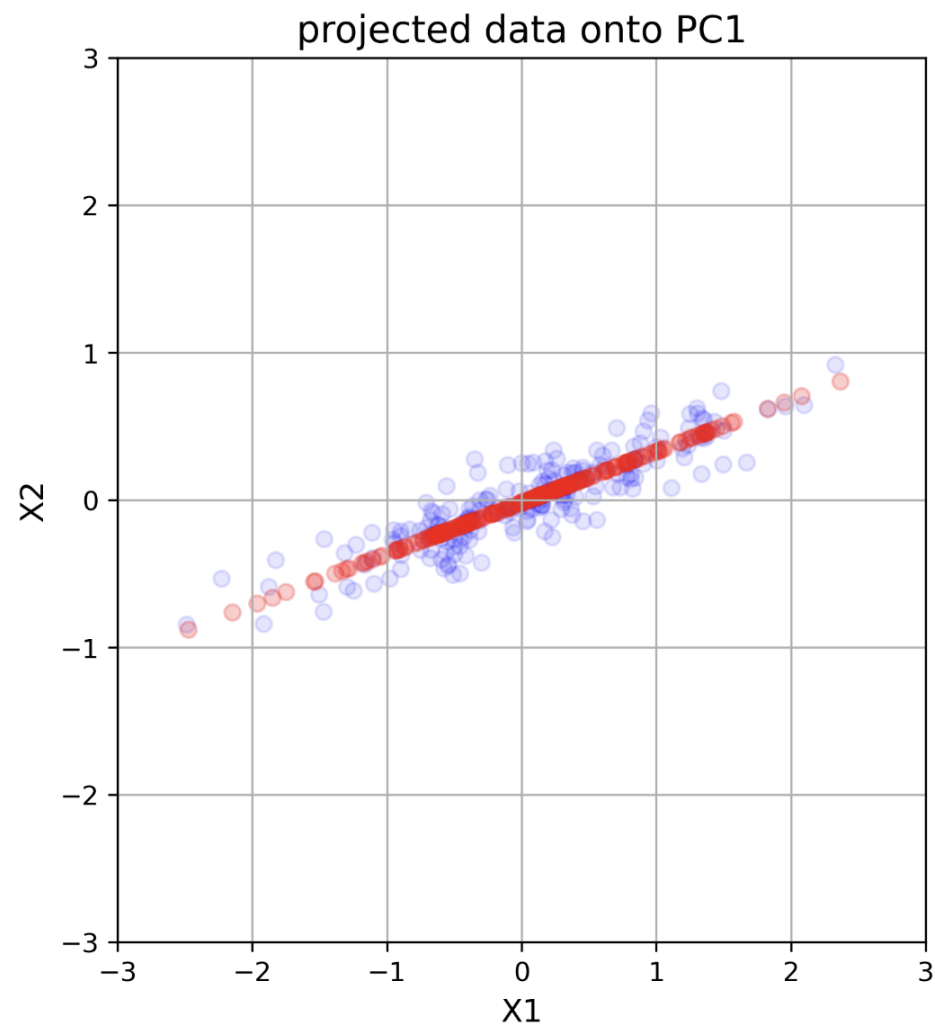
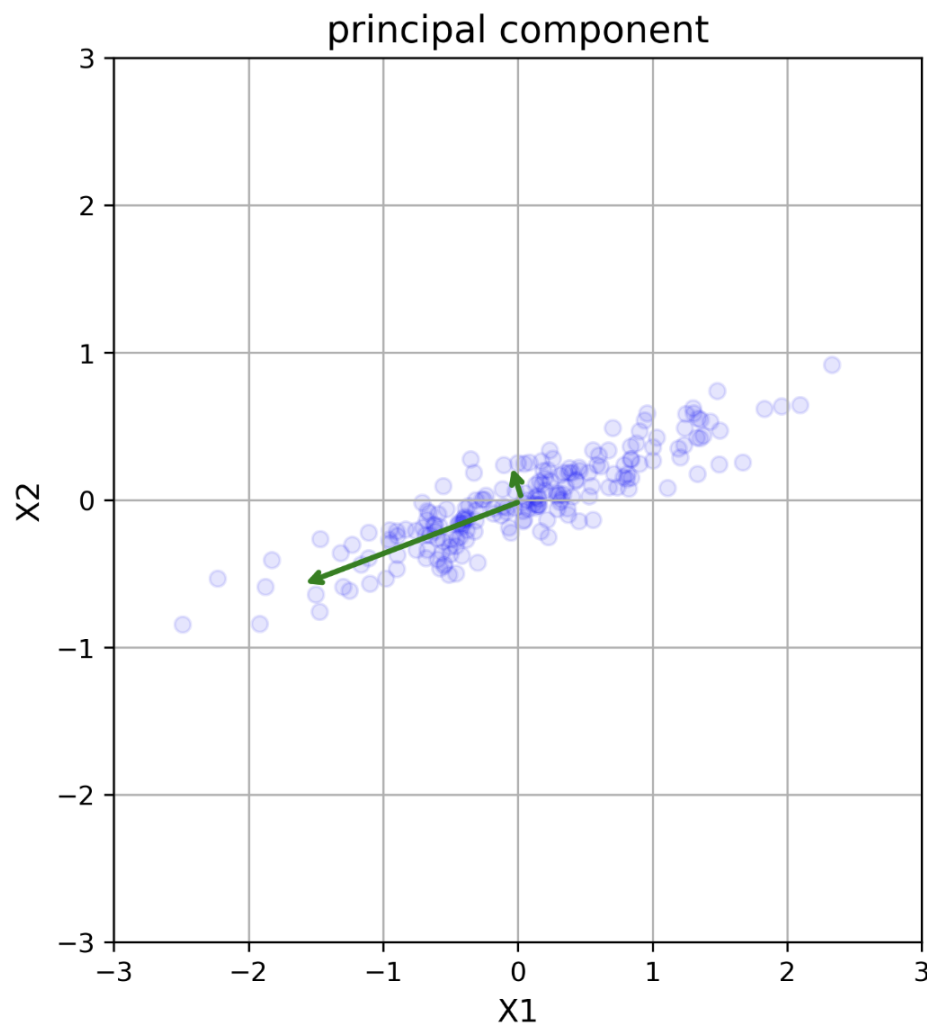
```
print("transformed shape:", A5D_pca5D.shape) transformed shape: (7, 5)
```

```
#  $A_{reconstructed} = A_{pca} \times v^T + \bar{A}$ 
```

```
A2D_new1D = pca1D.inverse_transform(A2D_pca1D)
```

```
A5D_new5D = pca5D.inverse_transform(A5D_pca5D)
```

PCA – `sklearn.decomposition.pca`



PCA – sklearn.decomposition.pca

convert to DataFrame

```
A5D_new5D = pd.DataFrame.from_records(A5D_pca5D,  
                                     columns=[f"PC{i+1}" for i in range(A5D_new5D.shape[1])])
```

A5D_new5D	PC1	PC2	PC3	PC4	PC5
0	1.696429	-0.411885	1.282764	2.246521	1.243323
1	-2.084069	1.617905	0.122180	-0.281795	-0.718424
2	3.261464	3.613608	-0.191616	-1.351178	0.764951
3	-4.207169	-2.286135	-2.229844	-0.958474	1.087402
4	5.788023	-2.471383	-1.677120	-0.112444	-0.922710
5	-1.237965	-1.631645	3.888311	-0.961178	-0.436948
6	-3.216714	1.569535	-1.194676	1.418547	-1.017594

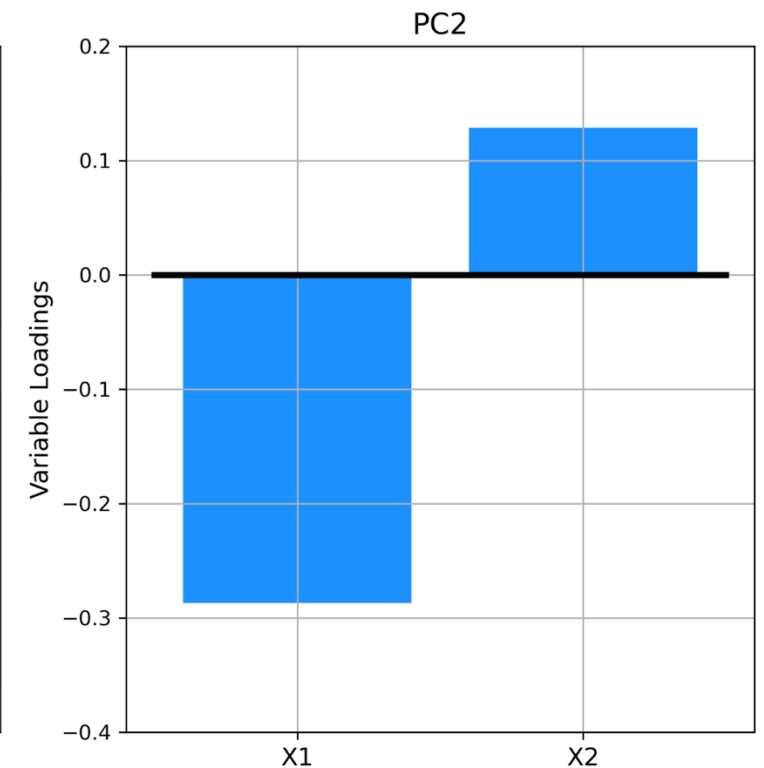
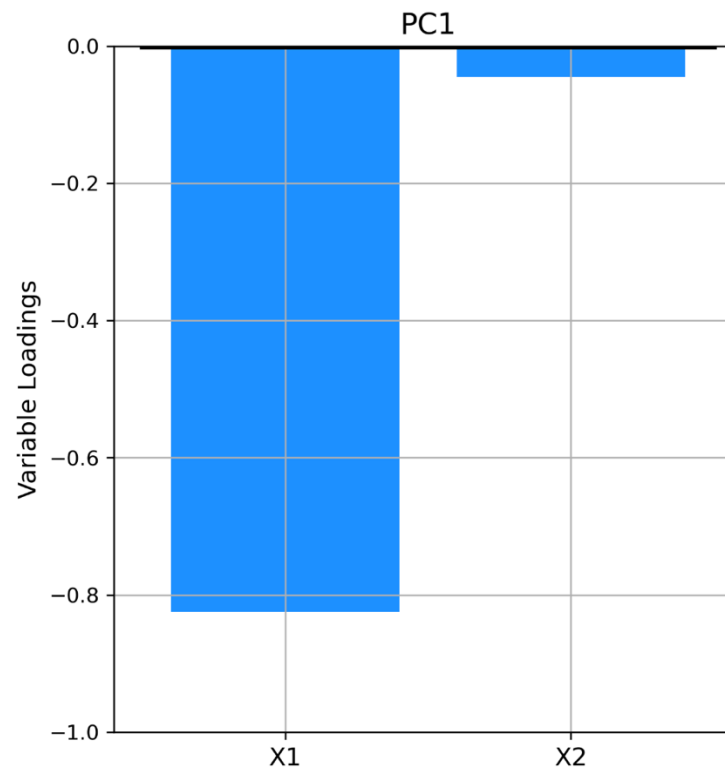
PCA – sklearn.decomposition.pca

variable loading calculation

```
variable_loading = pca2D.components_.T * np.sqrt(pca2D.explained_variance_)
```

```
print(variable_loading)
```

```
[[-0.82473153 -0.0446712 ]  
 [-0.28696587  0.12838372]]
```



PCA – Singular Value Decomposition

standardized matrix

A_centered = **A5D** - **np.mean**(**A5D**, axis=**0**)

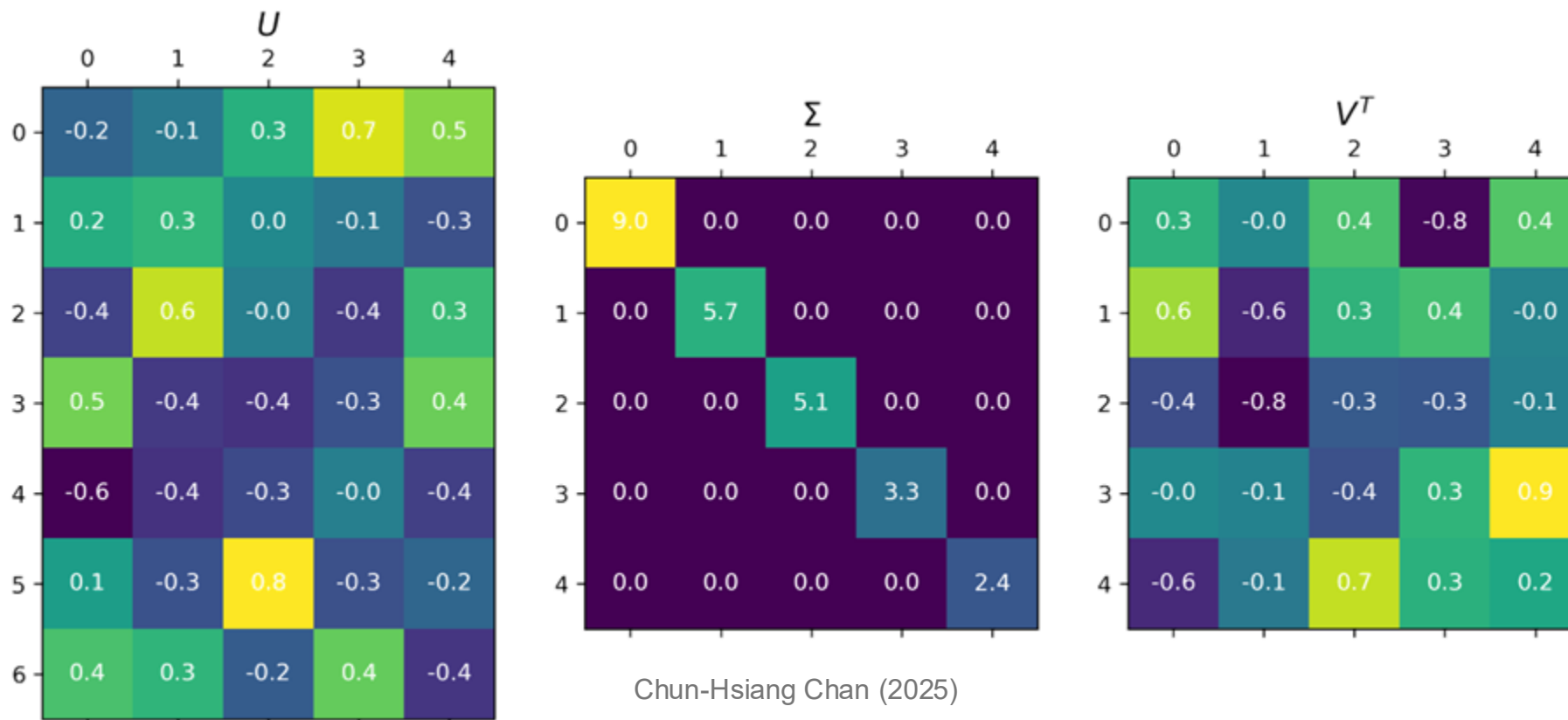
A_centered

```
array([[ -2.          ,  -1.          ,  -1.14285714,   1.71428571,   1.42857143],
       [  2.          ,  -1.          ,   0.85714286,  -1.28571429,   0.42857143],
       [  1.          ,  -2.          ,   0.85714286,   3.71428571,  -2.57142857],
       [  0.          ,   3.          ,   2.85714286,  -3.28571429,   1.42857143],
       [ -2.          ,   3.          ,  -3.14285714,   3.71428571,  -2.57142857],
       [ -2.          ,  -2.          ,  -1.14285714,  -3.28571429,  -0.57142857],
       [  3.          ,   0.          ,   0.85714286,  -1.28571429,   2.42857143]])
```

PCA – Singular Value Decomposition

compute SVD

U, S, Vt = `np.linalg.svd(A_centered, full_matrices=False)`



PCA – Singular Value Decomposition

proof $A = U\Sigma V^T$

- `print(np.round(A_centered, 3))`
- `print(np.round(U @ np.diag(S) @ Vt, 3))`




Diagram illustrating the result of the second code line: `print(np.round(U @ np.diag(S) @ Vt, 3))`. An orange line connects the code to the second matrix output.

<code>[[-2. -1. -1.143 1.714 1.429]</code>	<code>[[-2. -1. -1.143 1.714 1.429]</code>
<code>[2. -1. 0.857 -1.286 0.429]</code>	<code>[2. -1. 0.857 -1.286 0.429]</code>
<code>[1. -2. 0.857 3.714 -2.571]</code>	<code>[1. -2. 0.857 3.714 -2.571]</code>
<code>[0. 3. 2.857 -3.286 1.429]</code>	<code>[0. 3. 2.857 -3.286 1.429]</code>
<code>[-2. 3. -3.143 3.714 -2.571]</code>	<code>[-2. 3. -3.143 3.714 -2.571]</code>
<code>[-2. -2. -1.143 -3.286 -0.571]</code>	<code>[-2. -2. -1.143 -3.286 -0.571]</code>
<code>[3. 0. 0.857 -1.286 2.429]]</code>	<code>[3. 0. 0.857 -1.286 2.429]]</code>

PCA – Singular Value Decomposition

extract principal components (PC directions)

principal_components = Vt.T # Right singular vectors

principal_components

```
array([[ 2.84777513e-01,  6.37290518e-01, -4.04144877e-01,
        -4.55042776e-04, -5.91125431e-01],
       [-1.33828752e-02, -6.01836596e-01, -7.88343623e-01,
        -5.10904099e-02, -1.16265701e-01],
       [ 4.01328150e-01,  2.90984089e-01, -3.10944893e-01,
        -3.73205245e-01,  7.19927145e-01],
       [-7.56452377e-01,  3.81084174e-01, -3.38567133e-01,
        3.01380387e-01,  2.77663127e-01],
       [ 4.30625340e-01, -4.19120474e-02, -6.21837647e-02,
        8.75943646e-01,  2.04110524e-01]])
```

PCA – Singular Value Decomposition

compute projected data (Principal Component Scores)

A_pca = U @ np.diag(S)

print(A_pca)

```
[[-1.69642934 -0.41188488  1.28276422  2.24652093  1.24332308]
 [ 2.0840688   1.61790489  0.12218009 -0.28179454 -0.71842426]
 [-3.26146374  3.61360798 -0.19161578 -1.35117808  0.76495093]
 [ 4.20716867 -2.28613474 -2.22984393 -0.95847371  1.08740236]
 [-5.78802326 -2.4713829  -1.67711969 -0.11244403 -0.92270986]
 [ 1.23796474 -1.63164506  3.88831104 -0.96117788 -0.4369479 ]
 [ 3.21671412  1.56953472 -1.19467594  1.4185473  -1.01759434]]
```

equivalent to A_centered @ principal_components

A_pca1 = A_centered @ Vt.T

print(A_pca1)

```
[[-1.69642934 -0.41188488  1.28276422  2.24652093  1.24332308]
 [ 2.0840688   1.61790489  0.12218009 -0.28179454 -0.71842426]
 [-3.26146374  3.61360798 -0.19161578 -1.35117808  0.76495093]
 [ 4.20716867 -2.28613474 -2.22984393 -0.95847371  1.08740236]
 [-5.78802326 -2.4713829  -1.67711969 -0.11244403 -0.92270986]
 [ 1.23796474 -1.63164506  3.88831104 -0.96117788 -0.4369479 ]
 [ 3.21671412  1.56953472 -1.19467594  1.4185473  -1.01759434]]
```

PCA – Singular Value Decomposition

```
# compute Variance Explained = S2/(N-1)
variance_explained = (S**2) / (A5D.shape[0] - 1)
print(variance_explained)
[13.48994148  5.38421307  4.33804606  1.80323359  0.98456579]
# convert to DataFrame for better readability
A_pca_df = pd.DataFrame(A_pca, columns=[f"PC{i+1}" for i
                                     in range(A_pca.shape[1])])
```

A_pca_df

	PC1	PC2	PC3	PC4	PC5
0	-1.696429	-0.411885	1.282764	2.246521	1.243323
1	2.084069	1.617905	0.122180	-0.281795	-0.718424
2	-3.261464	3.613608	-0.191616	-1.351178	0.764951
3	4.207169	-2.286135	-2.229844	-0.958474	1.087402
4	-5.788023	-2.471383	-1.677120	-0.112444	-0.922710
5	1.237965	-1.631645	3.888311	-0.961178	-0.436948
6	3.216714	1.569535	-1.194676	1.418547	-1.017594

PCA – Eigen Decomposition

centered matrix

A_centered = **A5D** - **np.mean**(**A5D**, axis=**0**)

A_centered `array([[-2. , -1. , -1.14285714, 1.71428571, 1.42857143],
 [2. , -1. , 0.85714286, -1.28571429, 0.42857143],
 [1. , -2. , 0.85714286, 3.71428571, -2.57142857],
 [0. , 3. , 2.85714286, -3.28571429, 1.42857143],
 [-2. , 3. , -3.14285714, 3.71428571, -2.57142857],
 [-2. , -2. , -1.14285714, -3.28571429, -0.57142857],
 [3. , 0. , 0.85714286, -1.28571429, 2.42857143]])`

compute the covariance Matrix

S = **np.cov**(**A_centered**, rowvar=**False**)

S `array([[4.33333333e+00, -6.66666667e-01, 2.66666667e+00,
 -1.16666667e+00, 1.50000000e+00],
 [-6.66666667e-01, 4.66666667e+00, 7.40148683e-17,
 -1.48029737e-16, 1.66666667e-01],
 [2.66666667e+00, 7.40148683e-17, 3.80952381e+00,
 -3.04761905e+00, 1.90476190e+00],
 [-1.16666667e+00, -1.48029737e-16, -3.04761905e+00,
 9.23809524e+00, -3.85714286e+00],
 [1.50000000e+00, 1.66666667e-01, 1.90476190e+00,
 -3.85714286e+00, 3.95238095e+00]])`

PCA – Eigen Decomposition

compute eigenvalues and eigenvectors

eigenvalues, eigenvectors = `np.linalg.eig(S)`

`print(eigenvalues)`

```
[13.48994148  5.38421307  4.33804606  0.98456579  1.80323359]
```

`print(eigenvectors)`

```
[[ -2.84777513e-01  6.37290518e-01 -4.04144877e-01  5.91125431e-01  
  -4.55042776e-04]  
 [ 1.33828752e-02 -6.01836596e-01 -7.88343623e-01  1.16265701e-01  
  -5.10904099e-02]  
 [-4.01328150e-01  2.90984089e-01 -3.10944893e-01 -7.19927145e-01  
  -3.73205245e-01]  
 [ 7.56452377e-01  3.81084174e-01 -3.38567133e-01 -2.77663127e-01  
   3.01380387e-01]  
 [-4.30625340e-01 -4.19120474e-02 -6.21837647e-02 -2.04110524e-01  
   8.75943646e-01]]
```

PCA – Eigen Decomposition

sort eigenvectors by largest eigenvalues

sorted_indices = np.argsort(eigenvalues)[::-1] # descending order

eigenvalues = eigenvalues[sorted_indices]

eigenvectors = eigenvectors[:, sorted_indices]

print(sorted_indices) [0 1 2 4 3]

print(eigenvalues) [13.48994148 5.38421307 4.33804606 1.80323359 0.98456579]

print(eigenvectors) [[-2.84777513e-01 6.37290518e-01 -4.04144877e-01 -4.55042776e-04
5.91125431e-01]
[1.33828752e-02 -6.01836596e-01 -7.88343623e-01 -5.10904099e-02
1.16265701e-01]
[-4.01328150e-01 2.90984089e-01 -3.10944893e-01 -3.73205245e-01
-7.19927145e-01]
[7.56452377e-01 3.81084174e-01 -3.38567133e-01 3.01380387e-01
-2.77663127e-01]
[-4.30625340e-01 -4.19120474e-02 -6.21837647e-02 8.75943646e-01
-2.04110524e-01]]

PCA – Eigen Decomposition

```
# compute principal component scores (project data)
```

```
# transform data onto PCA space
```

```
A_pca = A_centered @ eigenvectors
```

```
print(A_pca)
```

```
[[ 1.69642934 -0.41188488  1.28276422  2.24652093 -1.24332308]
 [-2.0840688   1.61790489  0.12218009 -0.28179454  0.71842426]
 [ 3.26146374  3.61360798 -0.19161578 -1.35117808 -0.76495093]
 [-4.20716867 -2.28613474 -2.22984393 -0.95847371 -1.08740236]
 [ 5.78802326 -2.4713829  -1.67711969 -0.11244403  0.92270986]
 [-1.23796474 -1.63164506  3.88831104 -0.96117788  0.4369479 ]
 [-3.21671412  1.56953472 -1.19467594  1.4185473  1.01759434]]
```

PCA – Eigen Decomposition

convert to DataFrame for better readability

```
A_pca_df = pd.DataFrame(A_pca, columns=[f"PC{i+1}" for i  
in range(A_pca.shape[1])])
```

A_pca_df	PC1	PC2	PC3	PC4	PC5
0	1.696429	-0.411885	1.282764	2.246521	-1.243323
1	-2.084069	1.617905	0.122180	-0.281795	0.718424
2	3.261464	3.613608	-0.191616	-1.351178	-0.764951
3	-4.207169	-2.286135	-2.229844	-0.958474	-1.087402
4	5.788023	-2.471383	-1.677120	-0.112444	0.922710
5	-1.237965	-1.631645	3.888311	-0.961178	0.436948
6	-3.216714	1.569535	-1.194676	1.418547	1.017594

PCA – Variable Loading

PCA :: variable loading

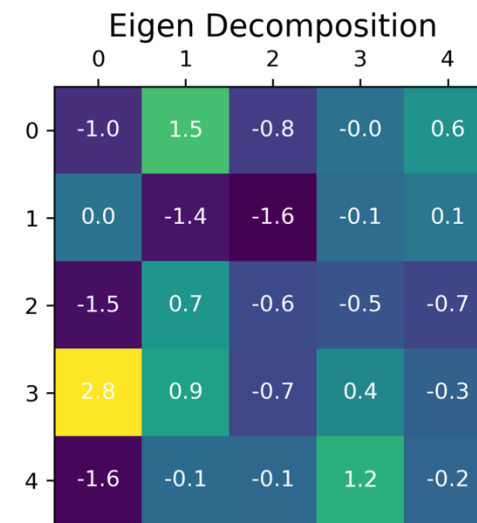
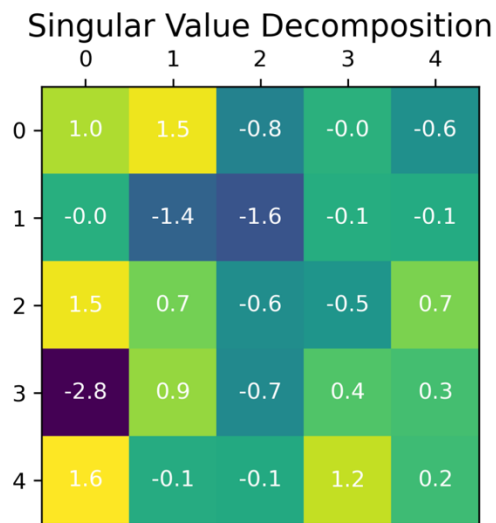
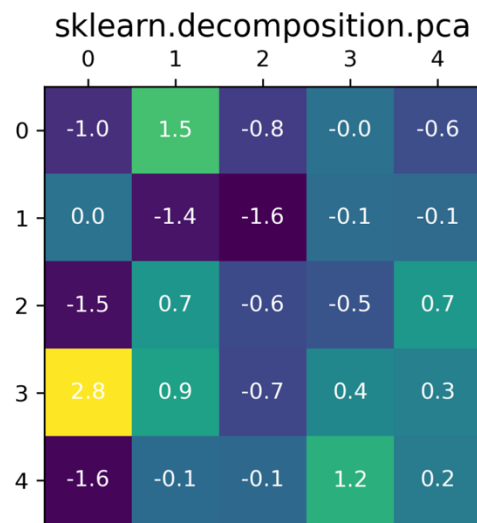
loading = pca5D.components_.T * `np.sqrt`(pca5D.explained_variance_)

SVD :: variable loading

loading1 = principal_components * `np.sqrt`(variance_explained)

Eigen :: variable loading

loading2 = eigenvectors * `np.sqrt`(eigenvalues)



The End

Thank you for your attention!

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